

Figure S1. Random-effects fit of model [2] (main text) to the raltegravir monotherapy data alone. The horizontal dashed line indicates the LOQ=400 copies/ml for the main assay used in the study. Data points below the dashed line were tested with a more sensitive assay with LOQ=50 copies/ml.

2 4 6 8

M11

M4

M8

5

4

3

2

M84

M88

M94

Viral load (log10)

5

4

3

2

M96

M1

M100

5

4

3

2

2

4

6

8

Time (days)

2

4

6

8

2 4 6 8

M15

M18

M7

5
4
3
2

M86

M91

M12

Viral load (log10)

5
4
3
2

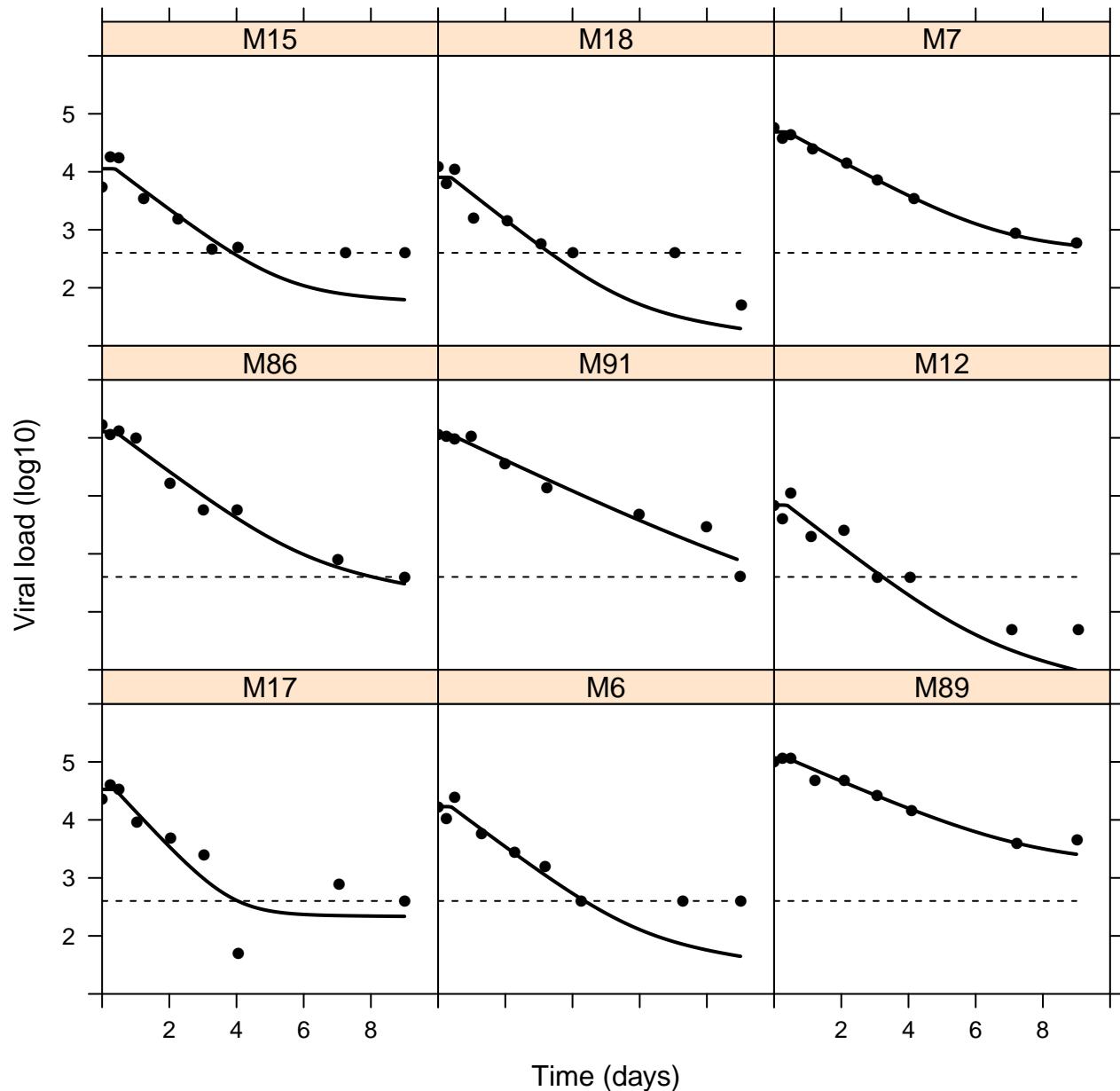
M17

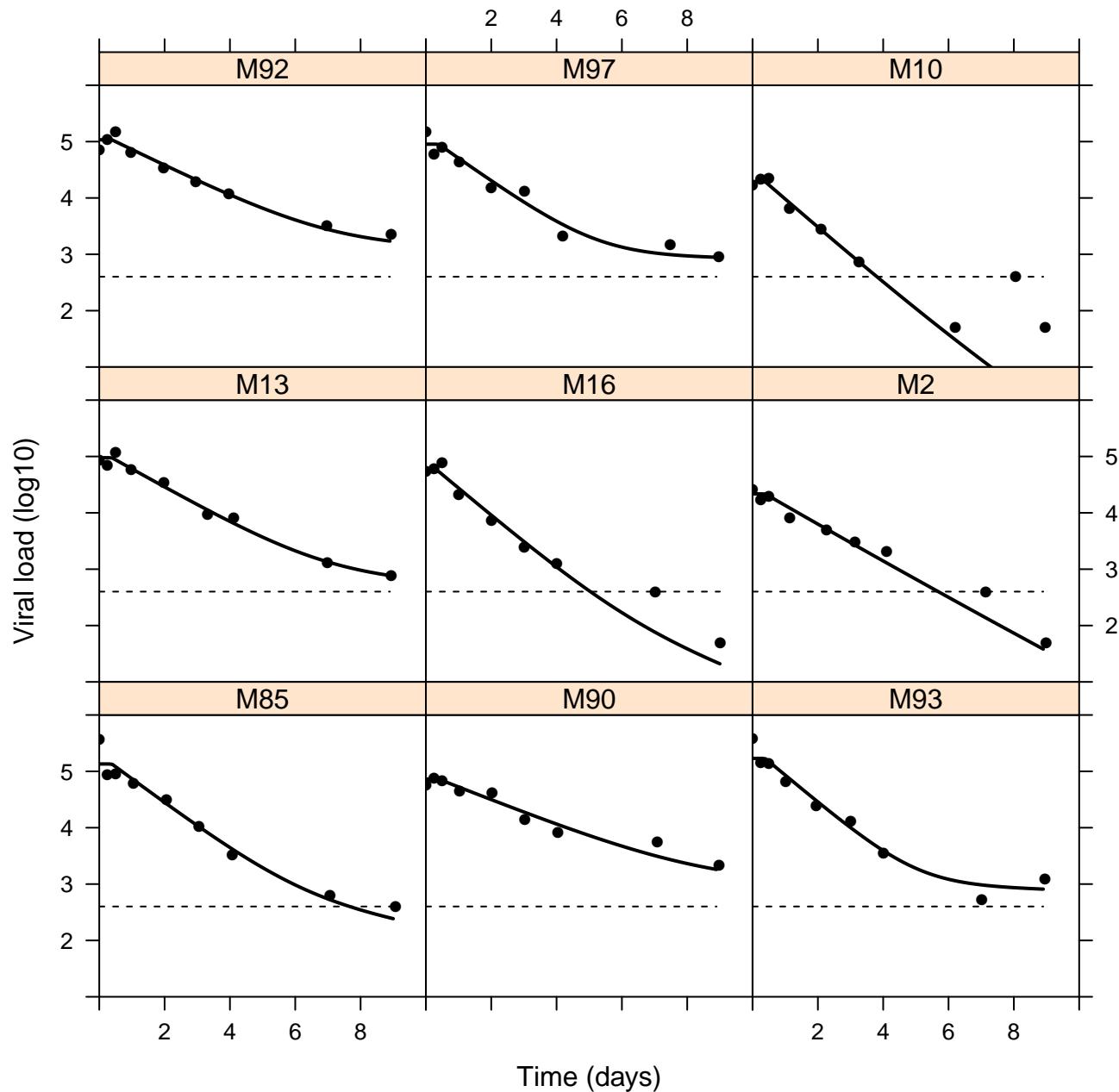
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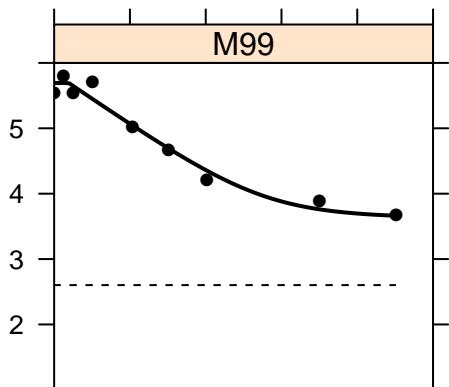
M89

5
4
3
2

Time (days)







Viral load (log₁₀)

Time (days)

Table S1. ESTIMATED PARAMETERS

	PID	V0	τ	δ_1	δ_2	ω	η	k	
Combination Tx	C1	4.79	0.47	0.08	0.72	0.938	0.95	2.6	
	C2	4.43	0.47	0.14	0.89	0.942	0.95	2.6	
	C5	4.59	0.47	0.14	1.31	0.945	0.95	2.6	
	C6	5.94	0.47	0.25	1.11	0.953	0.95	2.6	
	C13	4.64	0.47	0.25	0.67	0.932	0.95	2.6	
	C15	4.94	0.47	0.08	0.89	0.938	0.95	2.6	
	C16	4.42	0.47	0.13	0.81	0.938	0.95	2.6	
	C26	4.47	0.47	0.09	0.90	0.937	0.95	2.6	
	C27	4.42	0.47	0.12	0.97	0.944	0.95	2.6	
	C28	4.79	0.47	0.28	0.93	0.940	0.95	2.6	
	C32	4.95	0.47	0.18	0.78	0.940	0.95	2.6	
	Monotherapy	M1	3.89	0.47	0.19	1.14	0.997	0	2.6
		M2	4.32	0.47	0.52	0.79	0.997	0	2.6
M4		3.79	0.47	0.20	0.86	0.997	0	2.6	
M6		4.22	0.47	0.18	1.03	0.997	0	2.6	
M7		4.68	0.47	0.10	0.76	0.996	0	2.6	
M8		4.30	0.47	0.12	0.67	0.996	0	2.6	
M10		4.27	0.47	0.53	1.22	0.997	0	2.6	
M11		3.61	0.47	0.14	0.92	0.997	0	2.6	
M12		3.83	0.47	0.23	1.06	0.997	0	2.6	
M13		4.97	0.47	0.14	0.77	0.997	0	2.6	
M15		4.04	0.47	0.13	1.05	0.997	0	2.6	
M16		4.72	0.47	0.46	1.17	0.997	0	2.6	
M17		4.49	0.47	0.05	1.47	0.997	0	2.6	
M18		3.89	0.47	0.19	1.09	0.997	0	2.6	
M84		5.39	0.47	0.11	0.80	0.996	0	2.6	
M85		5.12	0.47	0.28	1.03	0.997	0	2.6	
M86		5.10	0.47	0.24	1.05	0.997	0	2.6	
M88		5.10	0.47	0.22	0.71	0.997	0	2.6	
M89		5.06	0.47	0.06	0.62	0.994	0	2.6	
M90		4.86	0.47	0.07	0.57	0.995	0	2.6	
M91		5.05	0.47	0.27	0.66	0.997	0	2.6	
M92		5.03	0.47	0.08	0.69	0.995	0	2.6	
M93		5.22	0.47	0.10	1.16	0.997	0	2.6	
M94		5.04	0.47	1.02	1.25	0.997	0	2.6	
M96		4.93	0.47	0.19	0.72	0.996	0	2.6	
M97		4.94	0.47	0.07	0.98	0.996	0	2.6	
M99		5.68	0.47	0.08	0.95	0.996	0	2.6	
M100		5.03	0.47	0.17	0.98	0.997	0	2.6	

Figure S2. Random-effects fit of model [2] (main text) to the monotherapy and combination therapy data together. Patients labeled C followed by a number were given combination therapy. The horizontal dashed line indicates the LOQ=400 copies/ml for the main assay used in the raltegravir monotherapy study. Data points below the dashed line were tested with a more sensitive assay with LOQ=50 copies/ml.

2 4 6 8 10

C1

C13

C15

6

5

4

3

2

C16

C2

C26

Viral load (log10)

6

5

4

3

2

C27

C28

C32

6

5

4

3

2

2

4

6

8

10

2

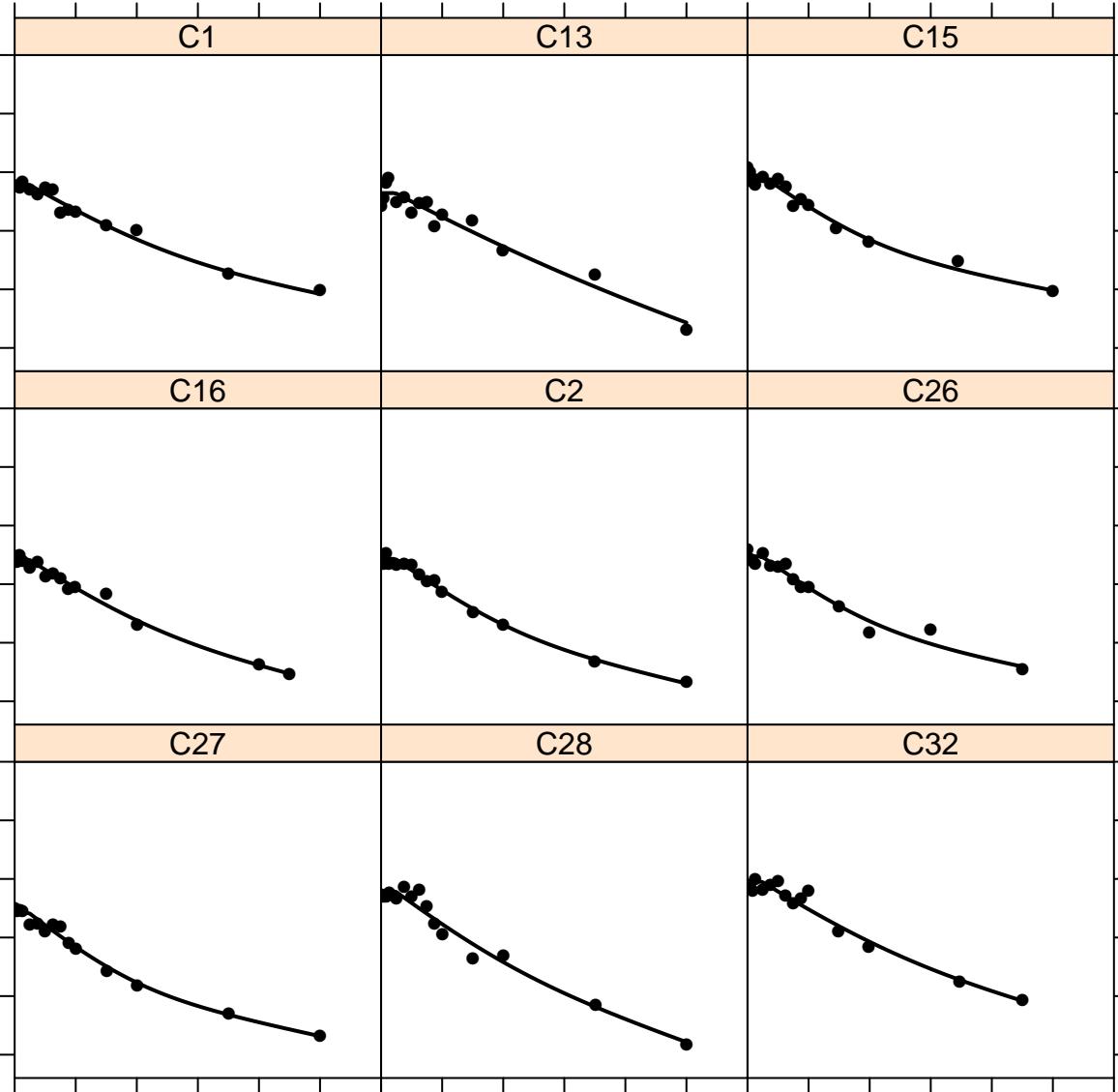
4

6

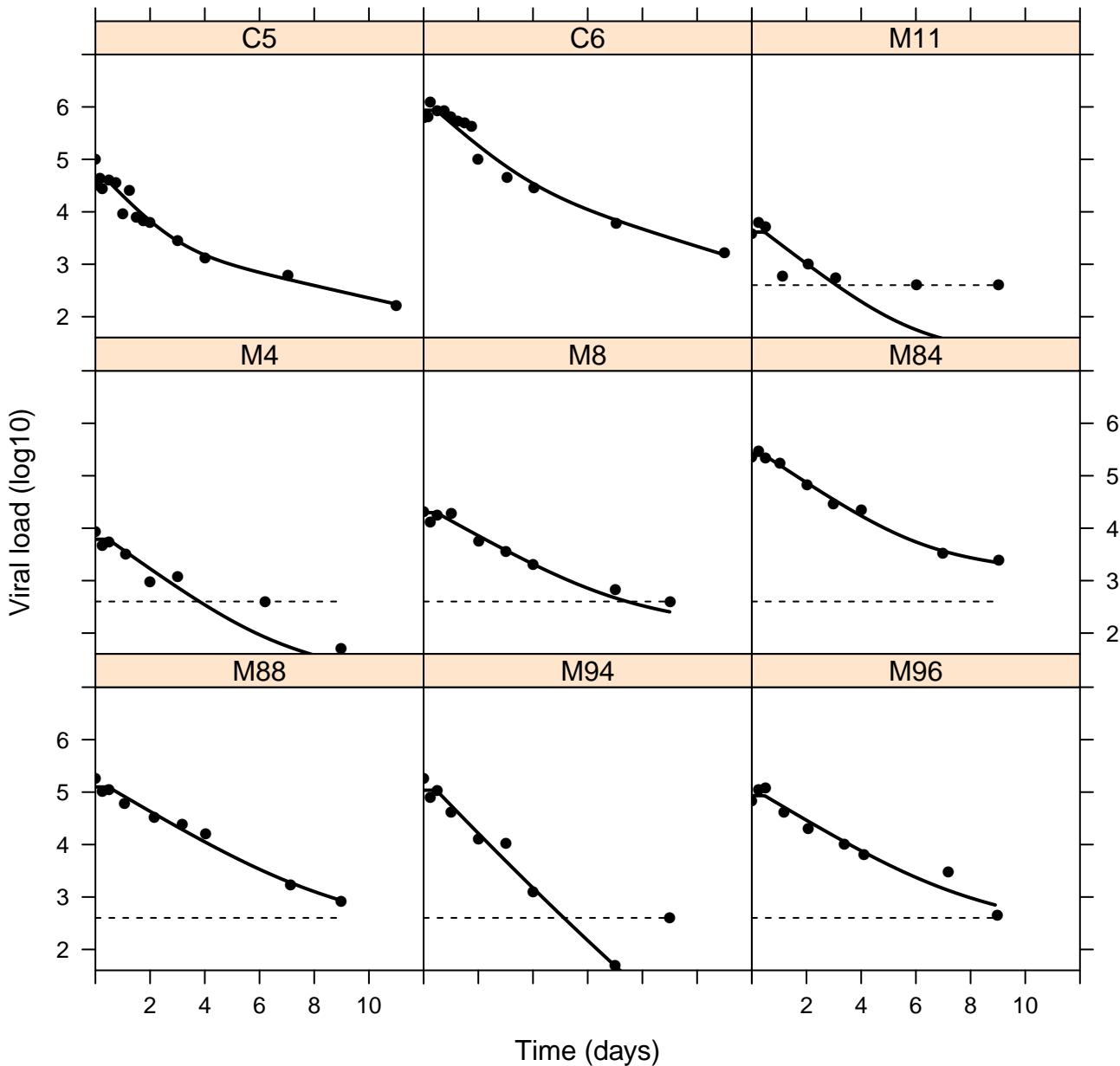
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10

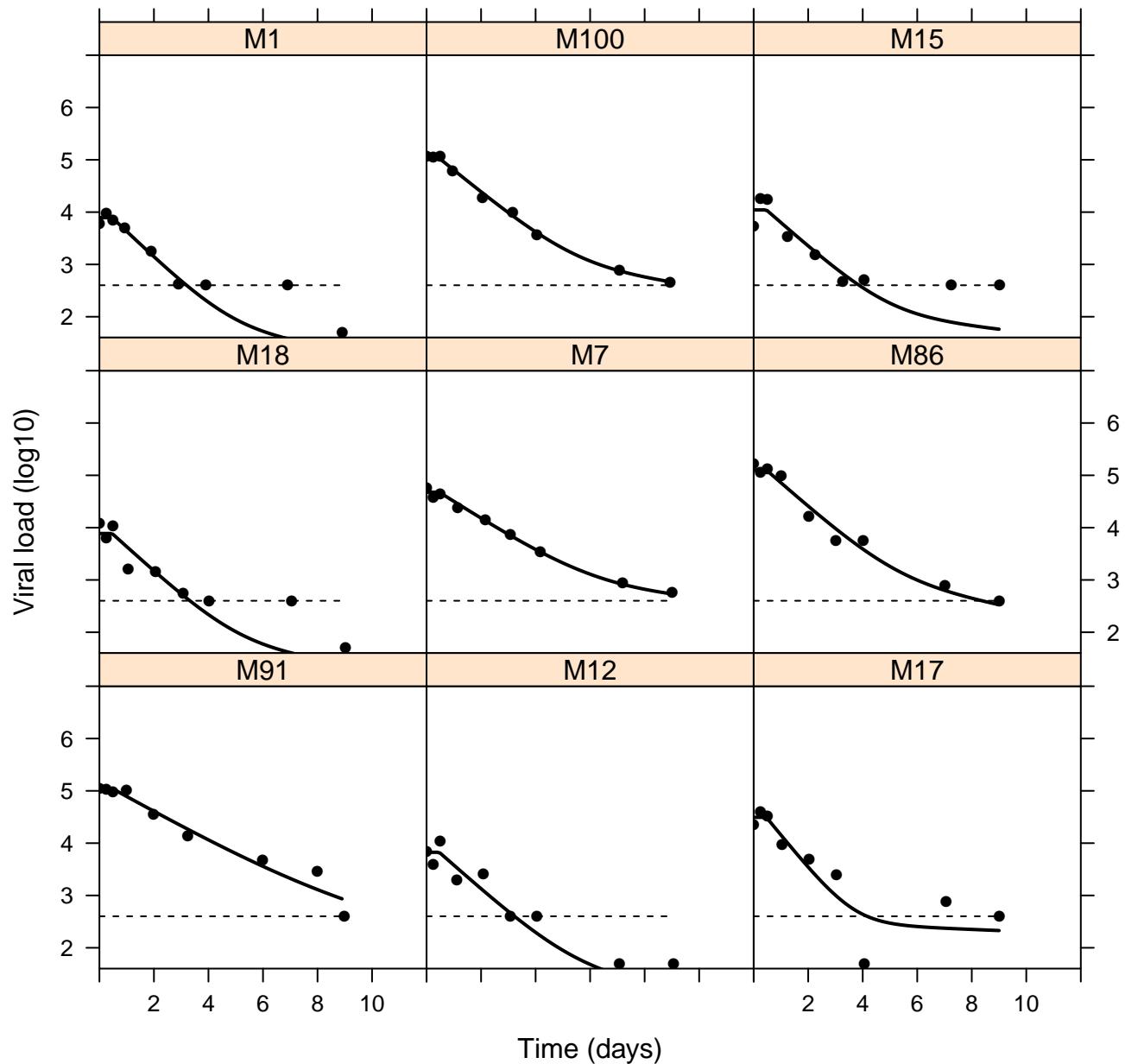
Time (days)



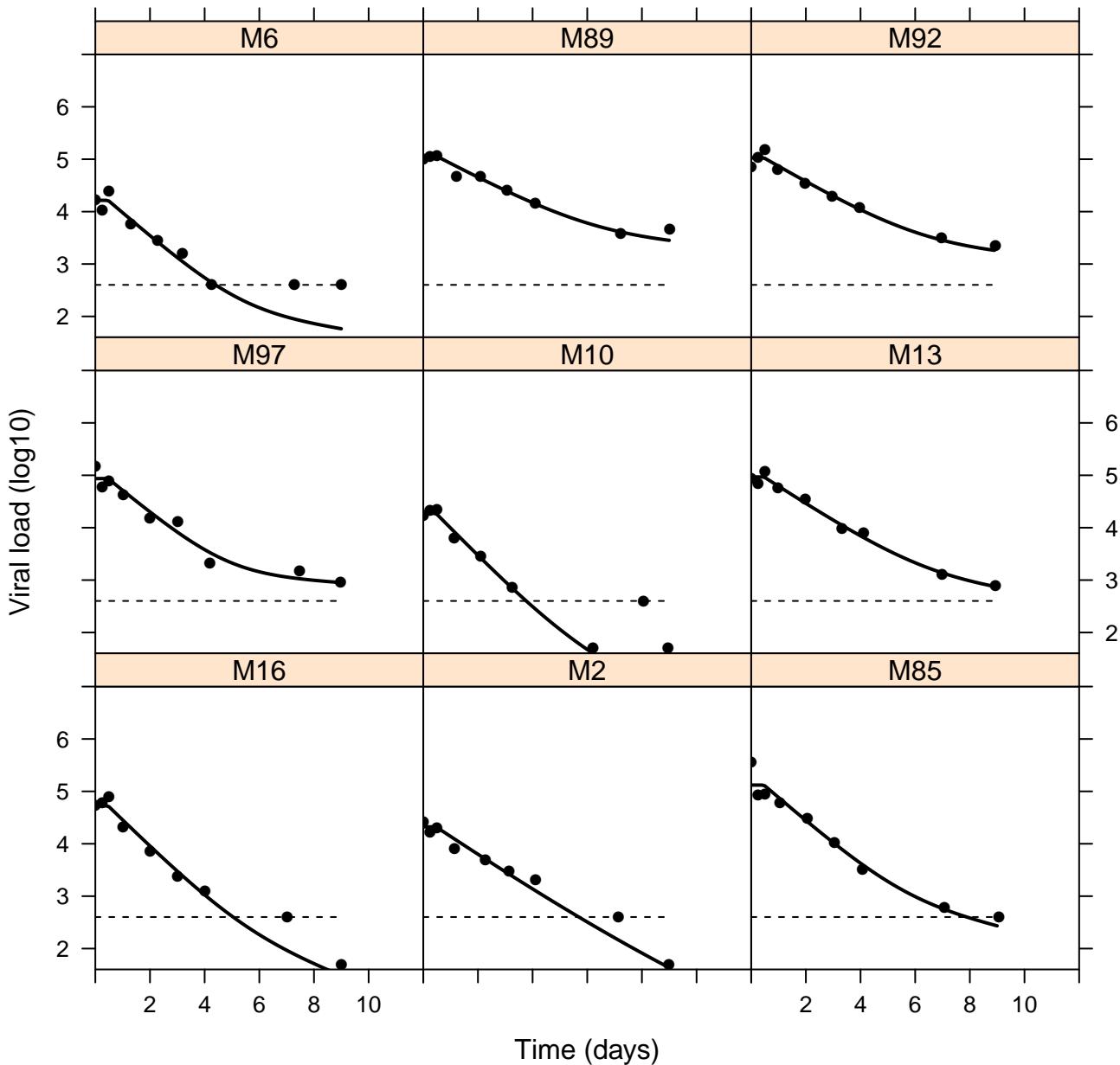
2 4 6 8 10



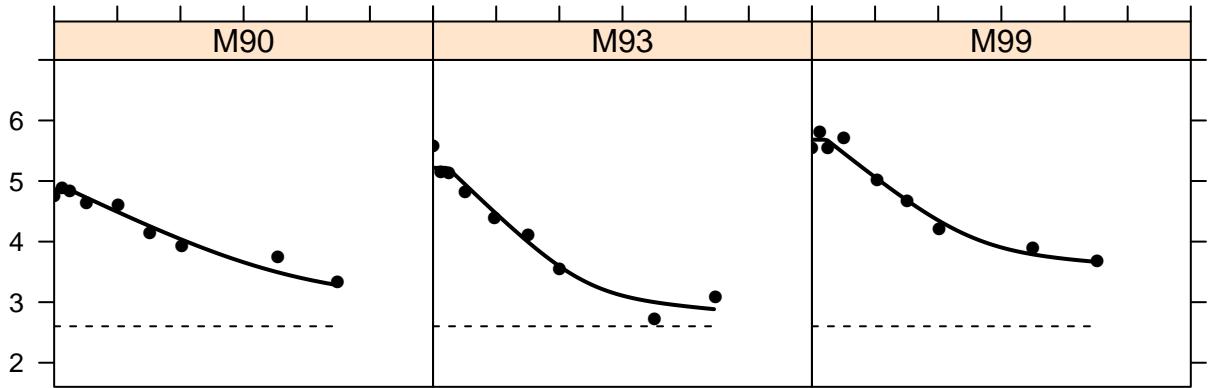
2 4 6 8 10



2 4 6 8 10



2 4 6 8 10



Viral load (log10)

Time (days)